

Errors Corrected by the STIC Systems Branch

Serial Number: 09/685,189

CRF Processing Date: 10/27/2002
 Edited by: _____
 Verified by: _____ (STIC staff)

ENTERED

RECEIVED
 MAR 9 2001
 TECH CENTER 1600/2800

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

***Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/685,189

DATE: 10/27/2000
 TIME: 08:34:57

Input Set : A:\ES.txt
 Output Set: N:\CRF3\10272000\I685189.raw

3 <110> APPLICANT: HEINRICHS, VOLKER
 4 CHEN, TEDDY
 5 PATTEN, PHILLIP A.
 7 <120> TITLE OF INVENTION: IFN-ALPHA HOMOLOGUES
 9 <130> FILE REFERENCE: 02-101510/0140.002
 OK 11 <140> CURRENT APPLICATION NUMBER: US/09/685,189
 12 <141> CURRENT FILING DATE: 2000-10-06
 14 <150> PRIOR APPLICATION NUMBER: 09/415,183
 15 <151> PRIOR FILING DATE: 1999-10-07
 17 <160> NUMBER OF SEQ ID NOS: 88
 19 <170> SOFTWARE: PatentIn Ver. 2.0

Does Not Comply
 Corrected Diskette Needed

ERRORED SEQUENCES

3185 <210> SEQ ID NO: 88
 3186 <211> LENGTH: 26
 3187 <212> TYPE: PRT
 3188 <213> ORGANISM: Artificial Sequence
 3190 <220> FEATURE:
 3191 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic amino acid
 3193 <400> SEQUENCE: 88
 3194 Trp Glu Leu Val Arg Ala Glu Ile Val Arg Ser Phe Ser Phe Ser Thr
 3195 1 5 10 15
 3197 Asn Leu Asn Lys Arg Leu Arg Lys Lys Glu
 3198 20 25

E--> 3205 1

delete at end of file

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/685,189

DATE: 10/27/2000

TIME: 08:34:59

Input Set : A:\ES.txt

Output Set: N:\CRF3\10272000\I685189.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2648 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2666 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2669 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:2672 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71
L:3205 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:88